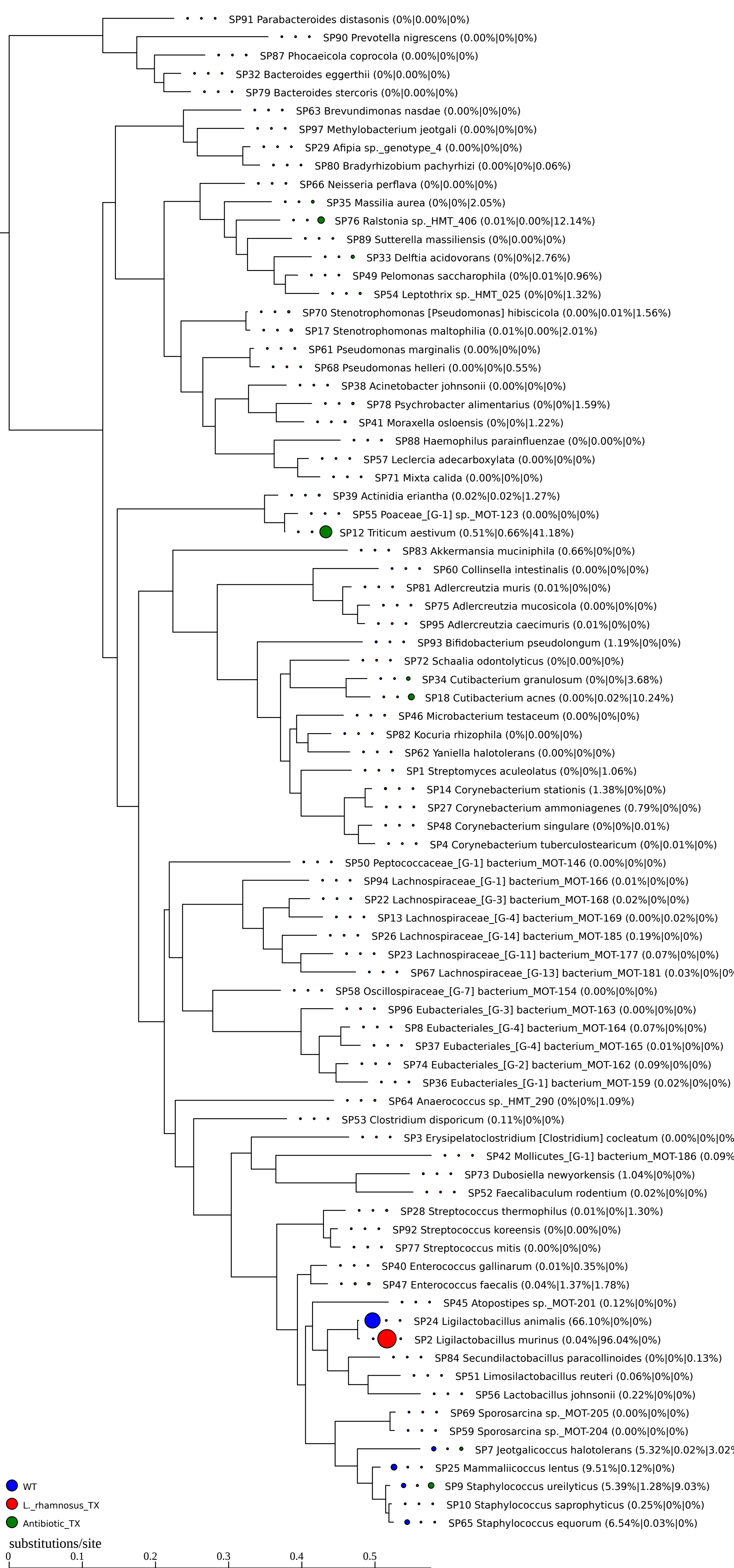
SP91 Parabacteroides distasonis (0%|0.00%|0%)
SP90 Prevotella nigrescens (0.00%|0%|0%)
SP87 Phocaeicola coprocola (0.00%|0%|0%)
SP32 Bacteroides eggerthii (0%|0.00%|0%)
SP79 Bacteroides stercoris (0%|0.00%|0%)
SP63 Brevundimonas nasdae (0.00%|0%|0%)
SP97 Methylobacterium jeotgali (0.00%|0%|0%)
SP29 Afipia sp._genotype_4 (0.00%|0%|0%)
SP80 Bradyrhizobium pachyrhizi (0.00%|0%|0.06%)
SP66 Neisseria perflava (0%|0.00%|0%)
SP35 Massilia aurea (0%|0%|2.05%)
SP76 Ralstonia sp._HMT_406 (0.01%|0.00%|12.14%)
SP89 Sutterella massiliensis (0%|0.00%|0%)
SP33 Delftia acidovorans (0%|0%|2.76%)
SP49 Pelomonas saccharophila (0%|0.01%|0.96%)
SP54 Leptothrix sp._HMT_025 (0%|0%|1.32%)
SP70 Stenotrophomonas [Pseudomonas] hibiscicola (0.00%|0.01%|1.56%)
SP17 Stenotrophomonas maltophilia (0.01%|0.00%|2.01%)
SP61 Pseudomonas marginalis (0.00%|0%|0%)
SP68 Pseudomonas helleri (0.00%|0%|0.55%)
SP38 Acinetobacter johnsonii (0.00%|0%|0%)
SP78 Psychrobacter alimentarius (0%|0%|1.59%)
SP41 Moraxella osloensis (0%|0%|1.22%)
SP88 Haemophilus parainfluenzae (0%|0.00%|0%)
SP57 Leclercia adecarboxylata (0.00%|0%|0%)
SP71 Mixta calida (0.00%|0%|0%)
SP39 Actinidia eriantha (0.02%|0.02%|1.27%)
SP55 Poaceae_[G-1] sp._MOT-123 (0.00%|0%|0%)
SP12 Triticum aestivum (0.51%|0.66%|41.18%)
SP83 Akkermansia muciniphila (0.66%|0%|0%)
SP60 Collinsella intestinalis (0.00%|0%|0%)
SP81 Adlercreutzia muris (0.01%|0%|0%)
SP75 Adlercreutzia mucosicola (0.00%|0%|0%)
SP95 Adlercreutzia caecimuris (0.01%|0%|0%)
SP93 Bifidobacterium pseudolongum (1.19%|0%|0%)
SP72 Schaalia odontolyticus (0%|0.00%|0%)
SP34 Cutibacterium granulosum (0%|0%|3.68%)
SP18 Cutibacterium acnes (0.00%|0.02%|10.24%)
SP46 Microbacterium testaceum (0.00%|0%|0%)
SP82 Kocuria rhizophila (0%|0.00%|0%)
SP62 Yaniella halotolerans (0.00%|0%|0%)
SP1 Streptomyces aculeolatus (0%|0%|1.06%)
SP14 Corynebacterium stationis (1.38%|0%|0%)
SP27 Corynebacterium ammoniagenes (0.79%|0%|0%)
SP48 Corynebacterium singulare (0%|0%|0.01%)
SP4 Corynebacterium tuberculostearicum (0%|0.01%|0%)
SP50 Peptococcaceae_[G-1] bacterium_MOT-146 (0.00%|0%|0%)
SP94 Lachnospiraceae_[G-1] bacterium_MOT-166 (0.01%|0%|0%)
SP22 Lachnospiraceae_[G-3] bacterium_MOT-168 (0.02%|0%|0%)
SP13 Lachnospiraceae_[G-4] bacterium_MOT-169 (0.00%|0.02%|0%)
SP26 Lachnospiraceae_[G-14] bacterium_MOT-185 (0.19%|0%|0%)
SP23 Lachnospiraceae_[G-11] bacterium_MOT-177 (0.07%|0%|0%)
SP67 Lachnospiraceae_[G-13] bacterium_MOT-181 (0.03%|0%|0%
SP58 Oscillospiraceae_[G-7] bacterium_MOT-154 (0.00%|0%|0%)
SP96 Eubacteriales_[G-3] bacterium_MOT-163 (0.00%|0%|0%)
SP8 Eubacteriales_[G-4] bacterium_MOT-164 (0.07%|0%|0%)
SP37 Eubacteriales_[G-4] bacterium_MOT-165 (0.01%|0%|0%)
SP74 Eubacteriales_[G-2] bacterium_MOT-162 (0.09%|0%|0%)
SP36 Eubacteriales_[G-1] bacterium_MOT-159 (0.02%|0%|0%)
SP64 Anaerococcus sp._HMT_290 (0%|0%|1.09%)
SP53 Clostridium disporicum (0.11%|0%|0%)
SP3 Erysipelatoclostridium [Clostridium] cocleatum (0.00%|0%|0%
SP42 Mollicutes_[G-1] bacterium_MOT-186 (0.09%
SP73 Dubosiella newyorkensis (1.04%|0%|0%)
SP52 Faecalibaculum rodentium (0.02%|0%|0%)
SP28 Streptococcus thermophilus (0.01%|0%|1.30%)
SP92 Streptococcus koreensis (0%|0.00%|0%)
SP77 Streptococcus mitis (0.00%|0%|0%)
SP40 Enterococcus gallinarum (0.01%|0.35%|0%)
SP47 Enterococcus faecalis (0.04%|1.37%|1.78%)
SP45 Atopostipes sp._MOT-201 (0.12%|0%|0%)
SP24 Ligilactobacillus animalis (66.10%|0%|0%)
SP2 Ligilactobacillus murinus (0.04%|96.04%|0%)
SP84 Secundilactobacillus paracollinoides (0%|0%|0.13%)
SP51 Limosilactobacillus reuteri (0.06%|0%|0%)
SP56 Lactobacillus johnsonii (0.22%|0%|0%)
SP69 Sporosarcina sp._MOT-205 (0.00%|0%|0%)
SP59 Sporosarcina sp._MOT-204 (0.00%|0%|0%)
SP7 Jeotgalicoccus halotolerans (5.32%|0.02%|3.02%
SP25 Mammaliicoccus lentus (9.51%|0.12%|0%)
SP9 Staphylococcus ureilyticus (5.39%|1.28%|9.03%)
SP10 Staphylococcus saprophyticus (0.25%|0%|0%)
SP65 Staphylococcus equorum (6.54%|0.03%|0%)
WT
L._rhamnosus_TX
Antibiotic_TX
substitutions/site
0
0.1
0.2
0.3
0.4
0.5